# EXHIBIT C



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## ExPASy Proteomics Server

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You are here: ExPASy CH > Tools > Primary structure analysis > Compute pl/Mw

### Compute pl/Mw

Theoretical pl/Mw (average) for the user-entered sequence:

6 <u>0</u>	5 <u>0</u>	4 <u>0</u>	3 <u>0</u>	2 <u>0</u>	1 <u>0</u>
VKPIYIVTPT	RNSRWSSPDN	AYPSVLGQTI	PRLLYAKSSP	FLGCLVKEIP	YFPPPAAKED
12 <u>0</u>	11 <u>0</u>	10 <u>0</u>	9 <u>0</u>	8 <u>0</u>	7 <u>0</u>
AVWVDGKART	FAVVDLSKMR	LSYRSLQPEE	VRSGGHDYEG	CGRRHGVRIR	Nashiqsavv
18 <u>0</u>	17 <u>0</u>	16 <u>0</u>	15 <u>0</u>	14 <u>0</u>	13 <u>0</u>
YGIAAENVID	GGGFGMLLRK	PTIGVGGNFA	TVLAFPAGVC	ELYYAIHKAS	AWVDSGAQLG
24 <u>0</u>	23 <u>0</u>	22 <u>0</u>	21 <u>0</u>	20 <u>0</u>	19 <u>0</u>
VFKIPKKASE	RLLPVPPTVT	SFGIVVAWKV	FWAVRGGGGE	HDKKSMGDDH	VKLVDANGTL
30 <u>0</u>	29 <u>0</u>	28 <u>0</u>	27 <u>0</u>	26 <u>0</u>	25 <u>0</u>
KFPELGMNAS	CQTLTPMMSS	ATFEAMYLGT	MIRVIAQGPT	VVAPQLPDDL	GAVDIINRWQ
36 <u>0</u>	35 <u>0</u>	34 <u>0</u>	33 <u>0</u>	32 <u>0</u>	31 <u>0</u>
RVWEQIFSTW	SDYVYEPFPK	NTFKPFAEYK	NIEDDLLNRN	IPFVHLGHRD	HCNEMSWIQS
42 <u>0</u>	41 <u>0</u>	40 <u>0</u>	39 <u>0</u>	38 <u>0</u>	37 <u>0</u>
AAPLSWSKEI	VNYWFAPGAG	RKGVLFNIQY	TPEWATPFPH	FDPYGATISA	LLKPGAGIMI
48 <u>0</u>	47 <u>0</u>	460	45 <u>0</u>	440	43 <u>0</u>
FQRLAITKGK	VWGQKYFKGN	NDVSTFSSGL	DIDLGRNEVV	NPRQAYANYR	YNYMEPYVSK
				50 <u>0</u> QSIPPLIKKY	490 VDPTDYFRNE

Theoretical pl/Mw: 9.05 / 55765.91

SERID NO: 2

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#### Compute pl/Mw

Theoretical pl/Mw (average) for the user-entered sequence:

VKPIYIVTPT 6 <u>0</u>	5 <u>0</u> RNSRWSSPDN	4 <u>0</u> AYPSVLGQTI	3 <u>0</u> PRLLYAKSSP	2 <u>0</u> FLGCLVKEIP	1 <u>0</u> YFPPPAAKED
12 <u>0</u> AVWVDGKART	11 <u>0</u> FAVVDLSKMR	10 <u>0</u> LSYRSLQPEE	9 <u>0</u> VRSGGHDYEG	8 <u>0</u> CGRRHGVRIR	7 <u>0</u> NASHIQSAVV
					13 <u>0</u> AWVDSGAQLG
					19 <u>0</u> VKLVDANGTL
30 <u>0</u> KFPELGMNAS					
					31 <u>0</u> HCNEMSWIQS
42 <u>0</u> AAPLSWSKEI					
48 <u>0</u> FQRLAITKGK	47 <u>0</u> VWGQKYFKGN	460 NDVSTFSSGL	450 DIDLGRNEVV	440 NPRQAYANYR	43 <u>0</u> YNYMEPYVSK
-				50 <u>0</u> OSIPPLIKKY	490 VDPTDYFRNE

SEQIDNO:4

Theoretical pl/Mw: 8.89 / 55734.85

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## ExPASy Proteomics Server

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#### Compute pl/Mw

Theoretical pl/Mw: 9.19 / 55624.88

Theoretical pl/Mw (average) for the user-entered sequence:

1 <u>0</u> YFPPPAAKED	2 <u>0</u> FLGCLVKEIP	3 <u>0</u> PRLLYAKSSP	4 <u>0</u> AYPSVLGQTI	5 <u>0</u> RNSRWSSPDN	VKPLYIITPT
7 <u>0</u>	8 <u>0</u>	9 <u>0</u>	10 <u>0</u>	11 <u>0</u>	12 <u>0</u>
NVSHIQSAVV	CGRRHSVRIR	VRSGGHDYEG	LSYRSLQPET	FAVVDLNKMR	AVWVDGKART
13 <u>0</u>	14 <u>0</u>	15 <u>0</u>	16 <u>0</u>	17 <u>0</u>	18 <u>0</u>
AWVDSGAQLG	ELYYAIYKAS	PTLAFPAGVC	PTIGVGGNFA	GGGFGMLLRK	YGIAAENVID
19 <u>0</u>	20 <u>0</u>	21 <u>0</u>	22 <u>0</u>	23 <u>0</u>	24 <u>0</u>
VKLVDANGKL	HDKKSMGDDH	FWAVRGGGGE	SFGIVVAWQV	KLLPVPPTVT	IFKISKTVSE
25 <u>0</u>	26 <u>0</u>	27 <u>0</u>	28 <u>0</u>	29 <u>0</u>	30 <u>0</u>
GAVDIINKWQ	VVAPQLPADL	MIRIIAQGPK	ATFEAMYLGT	CKTLTPLMSS	KFPELGMNPS
	32 <u>0</u> IPFVHLGHRD				
	38 <u>0</u> FDPYGATISA				
430	44 <u>0</u>	450	46 <u>0</u>	47 <u>0</u>	48 <u>0</u>
YNYMEPYVSK	NPRQAYANYR	DIDLGRNEVV	NDVSTYASGK	VWGQKYFKGN	FERLAITKGK
49 <u>0</u> VDPTDYFRNE	50 <u>0</u> QSIPPLIKKY	٠.			

SEQIDNO: 6

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